>lcl|47247 unnamed protein product

Length=521

Score = 78.2 bits (191), Expect = 2e-19, Method: Compositional matrix adjust.

Identities = 55/170 (32%), Positives = 84/170 (49%), Gaps = 25/170 (15%)

Query 3 SLNCIVAVSQNMGIGKNGDLPWPPLRNEFRYFQRMTT------TSSVEGKQNLVIMGKKT 56

+ + +VAV + GIG +PW + + ++F+ +TT K+N V+MG+KT

Sbjct 22 AFSLVVAVDERGGIGDGRSIPWN-VPEDMKFFRDVTTKLRGKNVKPSPAKRNAVVMGRKT 80

Query 57 WFSIPEKNRPLKGRINLVLSRELK----------EPPQGAH-----FLSRSLDDALKLTE 101

W SIP K RPL GR+N+VLS L E + H ++ L+ AL+L

Sbjct 81 WDSIPPKFRPLPGRLNVVLSSTLTTQHLLDGLPDEEKRNLHADSIVAVNGGLEQALQLLA 140

Query 102 QPELANKVDMVWIVGGSSVYKEAMNHPG-HL--KLFVTRIMQDFESDTFF 148

P ++ V+ +GG SVY EA+ P HL ++ T I S + F

Sbjct 141 SPNYTPSIETVYCIGGGSVYAEALRPPCVHLLQAIYRTTIRASESSCSVF 190